



Mapping changes in forest diversity and disease in North American temperate forests.

J. Antonio Guzmán Q., Jeannine Cavender-Bares (PI),
Philip A. Townsend, Jesús N. Pinto-Ledezma,
Jennifer Juzwik, Gerard Sapés, Jonathan Knott.

guzman@umn.edu

Project goals

Project goals (summary):

1. Model species distributions using remote sensing data as input variables.
 - ✓ Pinto-Ledezma and Cavender-Bares (2021) *Scientific Reports*.
2. **Scale forest diversity detection to map communities using satellite imagery.**
3. **Identify and differentiate threats to oaks species.**
 - ✓ Sapés et al. (2022) *Remote Sensing of Environment*.



Jeannine
Cavender-Bares (PI)



Philip A.
Townsend



Jesús N.
Pinto-Ledezma



Gerard
Sapés



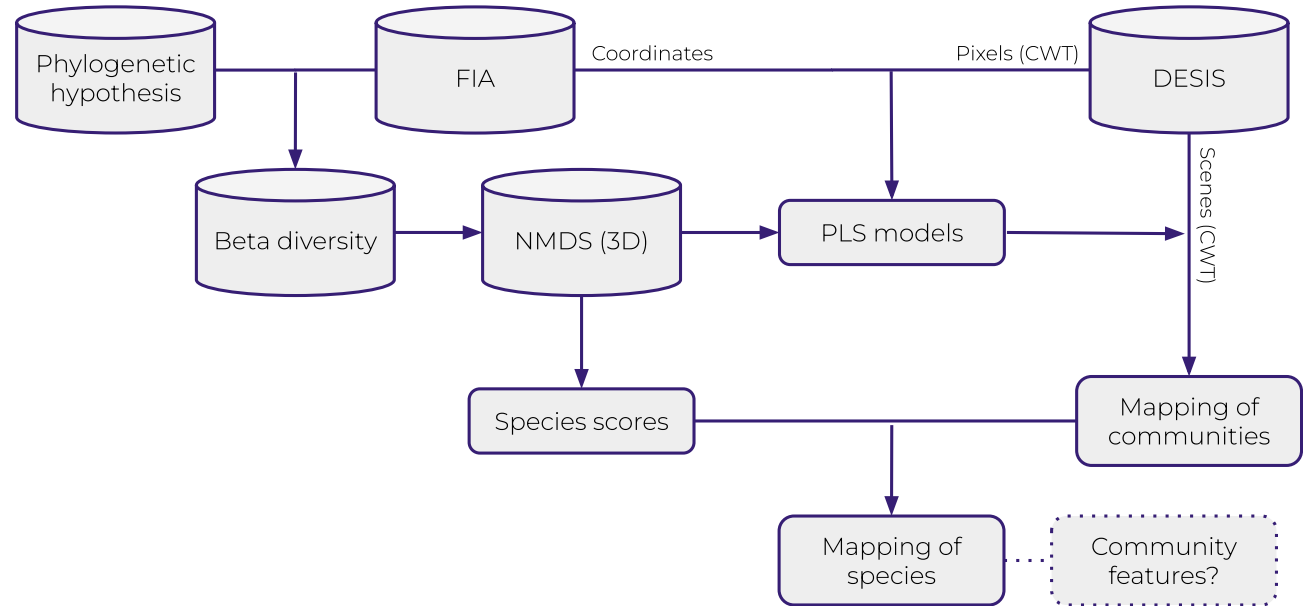
Jennifer
Juzwik



Jonathan
Knott

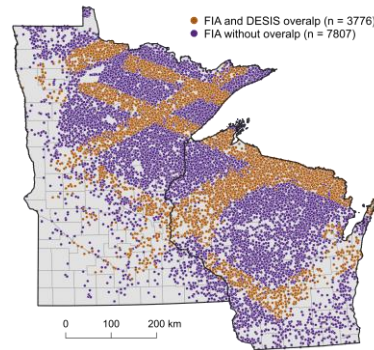
Mapping of forest communities

We are developing an approach to map forest communities by integrating forest inventories, phylogenetics, and space-borne spectroscopy



Schematic workflow for integrating phylogenetic beta diversity with spaceborne spectroscopy

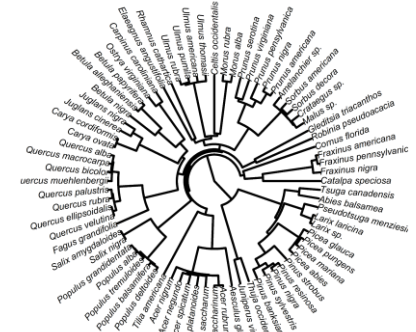
Mapping of forest communities



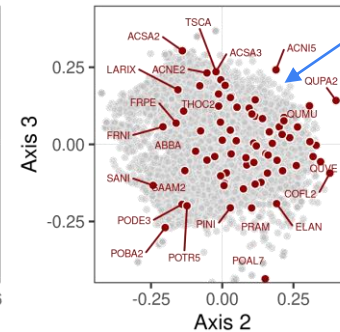
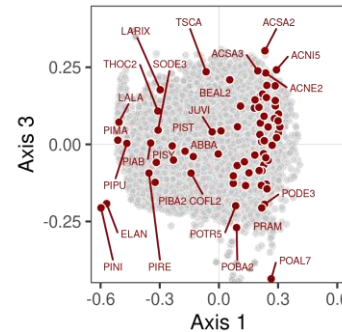
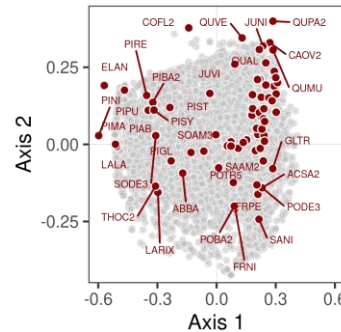
FIA plots



Phylogenetic beta diversity



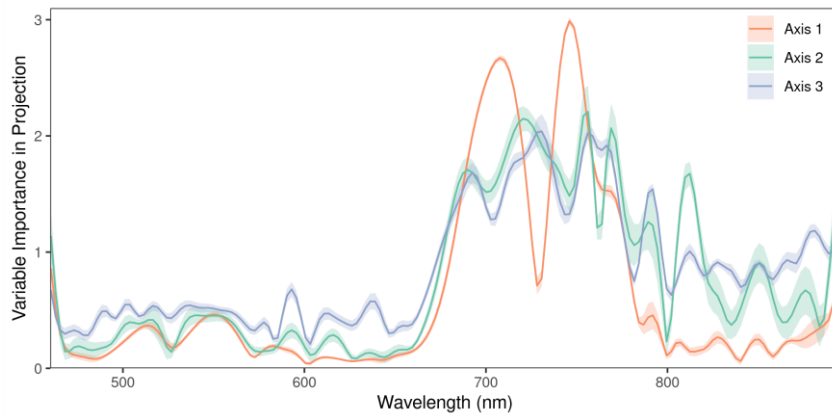
Phylogeny of trees
(72 species)



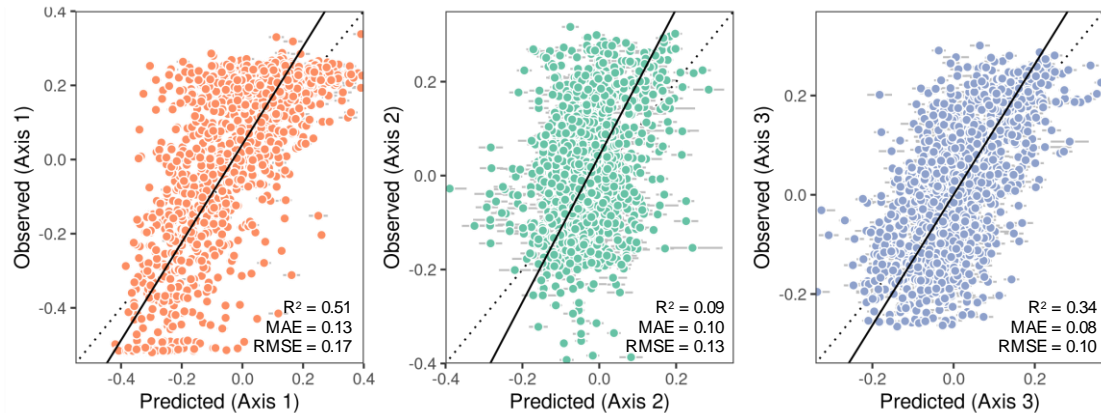
Non-metric Multi-Dimensional Scaling of phylogenetic beta diversity



Mapping of forest communities



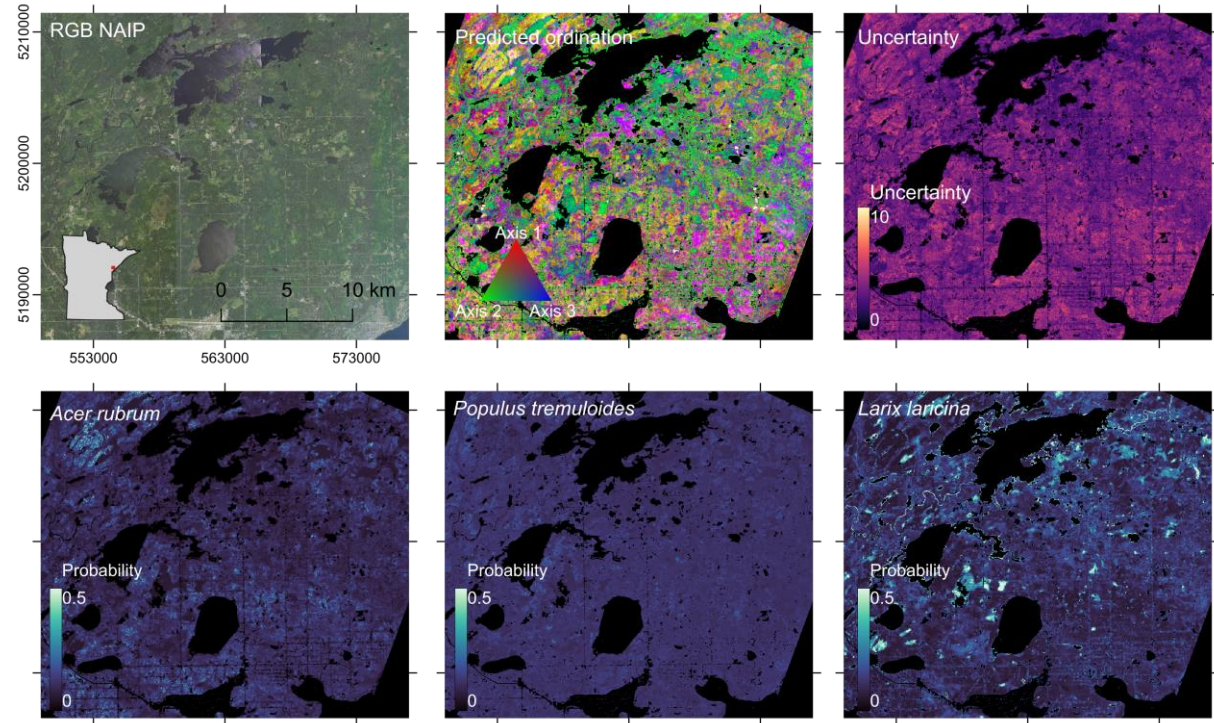
Variable importance of spectral bands that predict the NMDS axes.



Relationships between observed and predicted NMDS axes.

Prediction of the phylogenetic beta ordination axes for tree communities allows approximation of species occurrences

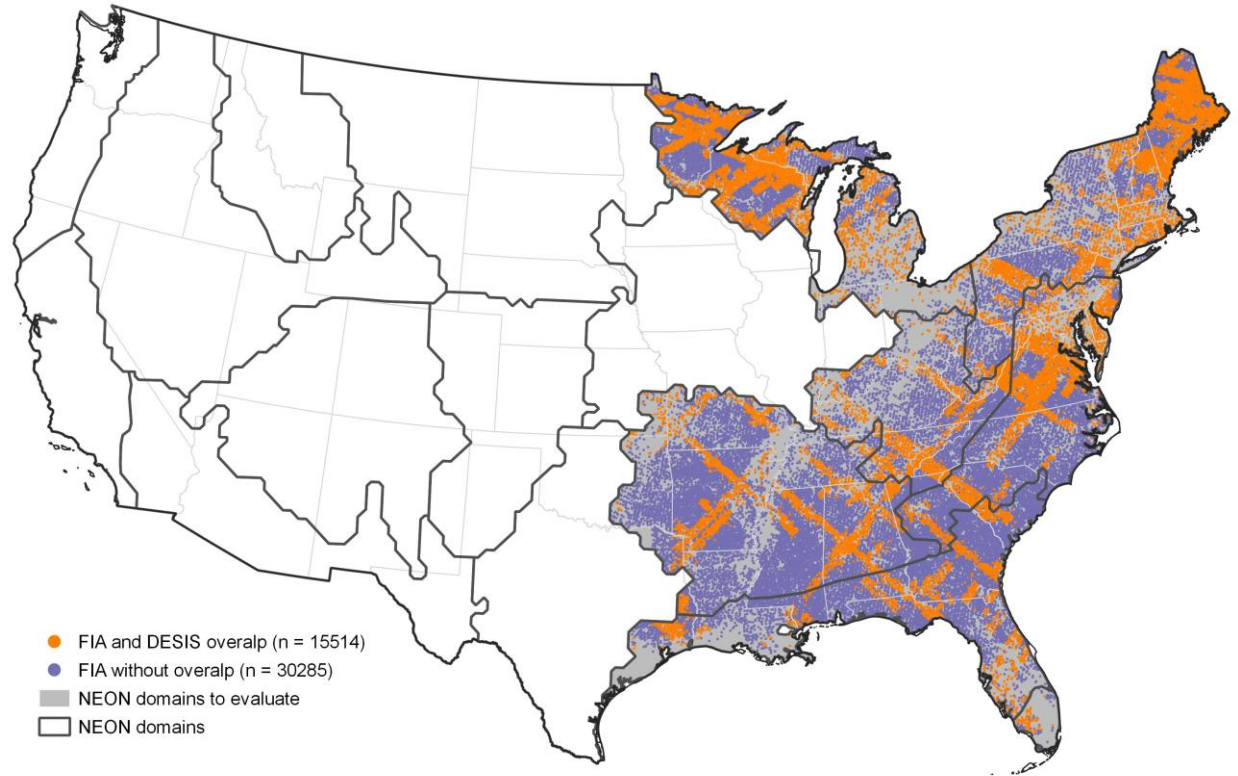
Mapping of forest communities



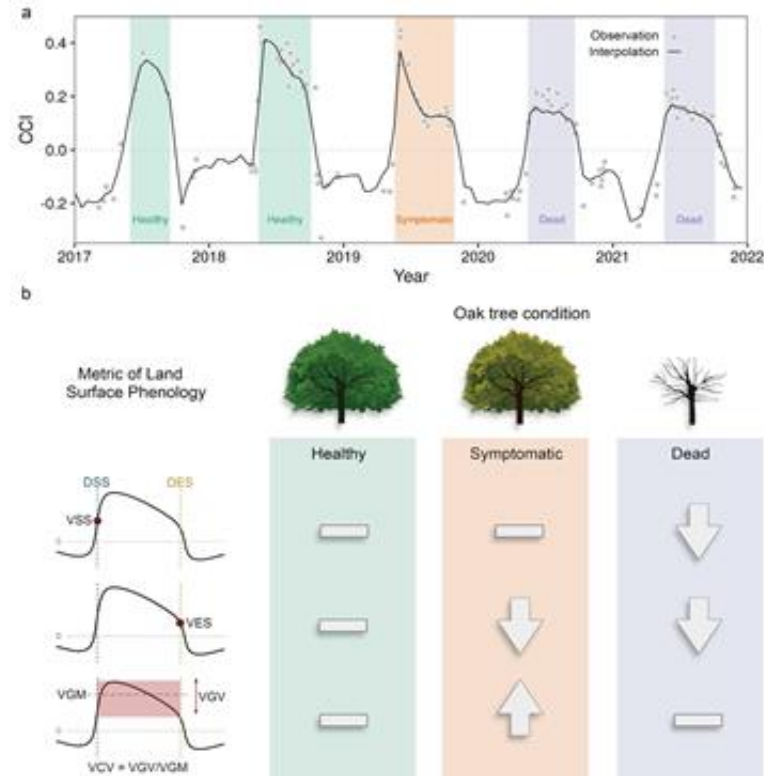
Mapping of the three community ordination axes (top middle panel) and the approximate species occurrences predicted from these models (bottom panels).

We are expanding our approach across Eastern US forests

Mapping
of forest
communities



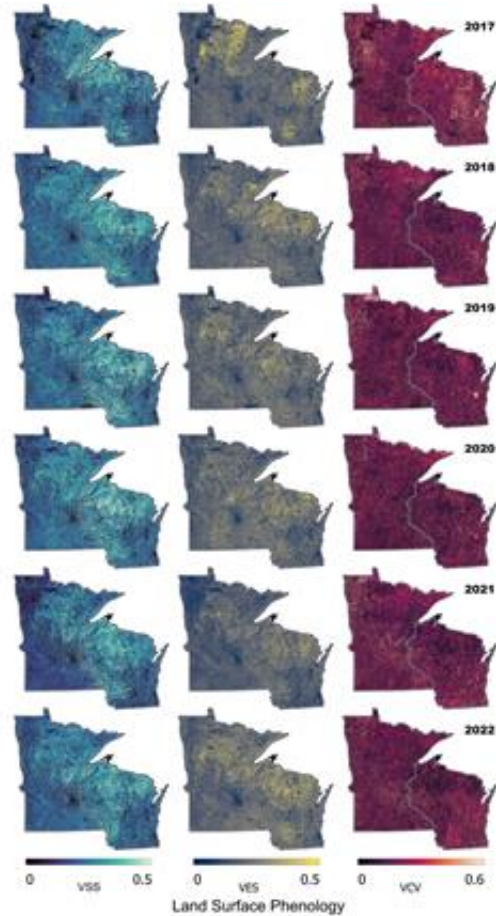
We have established that metrics of land surface phenology can be used as indicators of tree disease



Schematic representation of how metrics from land surface phenology can be used to infer the presence of oak wilt

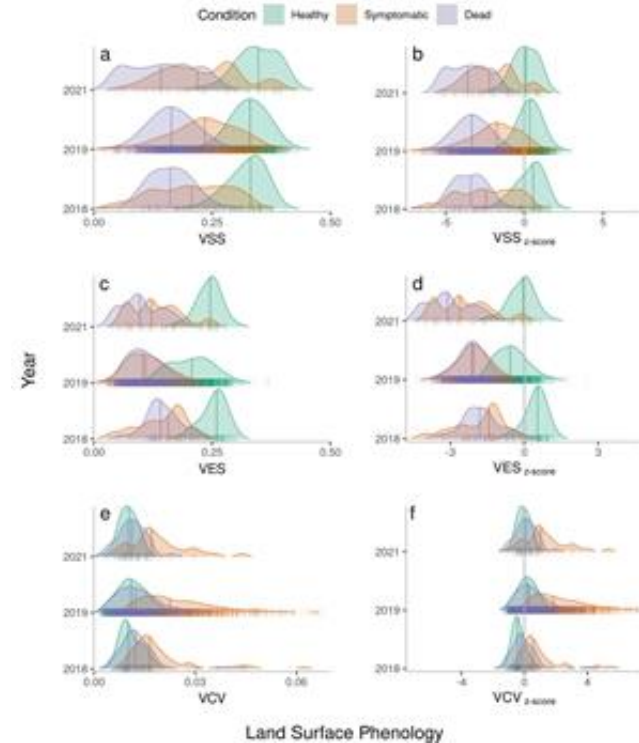
Mapping of
tree disease:
oak wilt

Mapping of tree disease: oak wilt



Land surface phenology across
Minnesota and Wisconsin derived from Sentinel-2

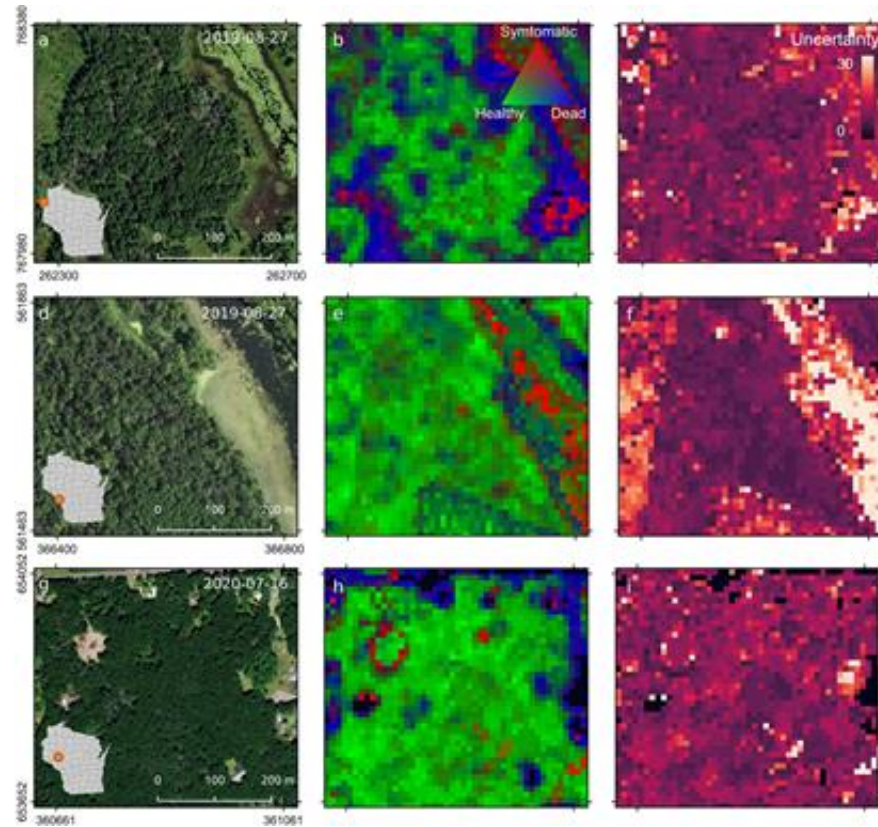
**We use land surface phenology to infer
oak wilt disease in oak trees**



Land surface phenology among oak
tree conditions

We created maps of the probability of oak wilt disease for both states

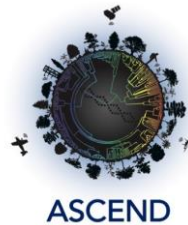
Mapping of
tree disease:
oak wilt



Oak wilt status for three sites in western Wisconsin. Panels represent NAIP RGB (left), probability maps (middle), and their uncertainty (right)

Thank you!

J. Antonio Guzmán Q., Ph.D
Department of Ecology, Evolution and Behavior
University of Minnesota, US
guzman@umn.edu



NASA Biodiversity Program (80NSSC21K1349)
Jeannine Cavender-Bares (PI)